

## SEQUENCE LISTING

<110> Keating, Mark T.  
Splawski, Igor

<120> MUTATIONS IN AND GENOMIC STRUCTURE OF HERG - A LONG QT  
SYNDROME GENE

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<150> US 09/735,995

<151> 2000-12-14

<150> US 09/226,012

<151> 1999-01-06

<150> US 09/122,847

<151> 1998-07-27

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<170> PatentIn Ver. 2.0

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 Ile Ala Asn Ala Arg Val Glu Asn Cys Ala Val Ile Tyr Cys Asn Asp

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Asp	Ile	Glu	Ala	Met	Arg	Ala	Gly	Val	Leu	Pro	Pro	Pro	Pro	Arg	His	

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caa Gln 335	atc Ile	acc Thr	ctc Leu	aac Asn	ttt Phe 340	gtg Val	gac Asp	ctc Leu	aag Lys	ggc Gly 345	gac Asp	ccc Pro	ttc Phe	ttg Leu	gct Ala 350	1116		
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gcc Ala	cgc Arg	tac Tyr	cac His	aca Thr 675	cag Gln	atg Met	ctg Leu	cgg Arg	gtg Val 680	cgg Arg	gag Glu	ttc Phe	atc Ile	cgc Arg 685	ttc Phe	2124
cac His	cag Gln	atc Ile	ccc Pro 690	aat Asn	ccc Pro	ctg Leu	cgc Arg	cag Gln 695	cgc Arg	ctc Leu	gag Glu	gag Glu	tac Tyr 700	ttc Phe	cag Gln	2172
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gac Asp	aca Thr	ctg Leu	gtg Val 770	cat His	gct Ala	ggg Gly	gac Asp	ctg Leu 775	ctc Leu	acc Thr	gcc Ala	ctg Leu	tac Tyr 780	ttc Phe	atc Ile	2412
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Met Thr Leu Val Pro Pro Ala Tyr Ser Ala Val Thr Thr Pro Gly Pro	1075	1080	1085	
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Gly Pro Thr Ser Thr Ser Pro Leu Leu Pro Val Ser Pro Leu Pro Thr	1090	1095	1100	
ctc acc ttg gac tcg ctt tct cag gtt tcc cag ttc atg gcg tgt gag				3420
Leu Thr Leu Asp Ser Leu Ser Gln Val Ser Gln Phe Met Ala Cys Glu	1105	1110	1115	
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Glu Leu Pro Pro Gly Ala Pro Glu Leu Pro Gln Glu Gly Pro Thr Arg	1120	1125	1130	
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Cys Glu Leu Cys Gly Tyr Ser Arg Ala Glu Val Met Gln Arg Pro Cys	50	55	60	
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Gln Ile Ala Gln Ala Leu Leu Gly Ala Glu Glu Arg Lys Val Glu Ile	85	90	95	

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Glu	Val	Val	Ser	His	Pro	Gly	Arg	Ile	Ala	Val	His	Tyr	Phe	Lys	Gly
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	690					695					700				
Trp	Ser	Tyr	Thr	Asn	Gly	Ile	Asp	Met	Asn	Ala	Val	Leu	Lys	Gly	Phe
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Pro	Glu	Cys	Leu	Gln	Ala	Asp	Ile	Cys	Leu	His	Leu	Asn	Arg	Ser	Leu
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 Lys Ser Asn Gly Asp Val Arg Ala Leu Thr Tyr Cys Asp Leu His Lys  
 820 825 830  
 Ile His Arg Asp Asp Leu Leu Glu Val Leu Asp Met Tyr Pro Glu Phe  
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 Ser Asp His Phe Trp Ser Ser Leu Glu Ile Thr Phe Asn Leu Arg Asp  
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 Thr Asn Met Ile Pro Gly Ser Pro Gly Ser Thr Glu Leu Glu Gly Gly  
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 945 950 955 960  
 Pro Arg Pro Pro Gly Glu Pro Pro Gly Gly Glu Pro Leu Met Glu Asp  
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 Cys Glu Lys Ser Ser Asp Thr Cys Asn Pro Leu Ser Gly Ala Phe Ser  
 980 985 990  
 Gly Val Ser Asn Ile Phe Ser Phe Trp Gly Asp Ser Arg Gly Arg Gln  
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 Thr Ser Thr Ser Pro Leu Leu Pro Val Ser Pro Leu Pro Thr Leu Thr  
 1090 1095 1100  
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 Pro Pro Gly Ala Pro Glu Leu Pro Gln Glu Gly Pro Thr Arg Arg Leu

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&lt;400&gt; 115

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